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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/101,825DATE: 01/14/2000
TIME: 00:09:20

INPUT SET: S34448.raw

This Raw Listing contains the General
Information Section and those Sequences
containing ERRORS.

SEQUENCE LISTING

1
2
3 (1) General Information:
4
5 (i) APPLICANT: GRONHOJ LARSEN, Christian
6 GESSER, Borbala
7
8 (ii) TITLE OF INVENTION: IMMUNOMODULATORS
9
10 (iii) NUMBER OF SEQUENCES: 23
11
12 (iv) CORRESPONDENCE ADDRESS:
13 (A) ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
14 (B) STREET: 624 Ninth Street N.W., Suite 300
15 (C) CITY: Washington
16 (D) STATE: D.C.
17 (E) COUNTRY: U.S.A.
18 (F) ZIP: 20001
19
20 (v) COMPUTER READABLE FORM:
21 (A) MEDIUM TYPE: Floppy disk
22 (B) COMPUTER: IBM PC compatible
23 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
24 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
25
26 (vi) CURRENT APPLICATION DATA:
27 (A) APPLICATION NUMBER: US 09/101,825
28 (B) FILING DATE: 17-JUL-1998
29
30 (vii) PRIOR APPLICATION DATA:
31 (A) APPLICATION NUMBER: PCT/DK97/00021
32 (B) FILING DATE: 16-JAN-1997
33
34 (vii) PRIOR APPLICATION DATA:
35 (A) APPLICATION NUMBER: PCT/DK96/00029
36 (B) FILING DATE: 18-JAN-1996
37
38 (viii) ATTORNEY/AGENT INFORMATION:
39 (A) NAME: COOPER, Iver P.
40 (B) REGISTRATION NUMBER: 28,005
41 (C) REFERENCE/DOCKET NUMBER: GRONHOJ-LARSEN=2
42
43 (ix) TELECOMMUNICATION INFORMATION:
44 (A) TELEPHONE: (202)628-5197
45 (B) TELEFAX: (202)737-3528

Does Not Comply
Corrected Diskette Needed

INPUT SET: S34448.raw

46
47

ERRORED SEQUENCES FOLLOW:

338 (2) INFORMATION FOR SEQ ID NO:19:
339
340 (i) SEQUENCE CHARACTERISTICS:
341 (A) LENGTH: 6 amino acids
342 (B) TYPE: amino acid
343 (C) STRANDEDNESS: single
344 (D) TOPOLOGY: linear
345
346 (ii) MOLECULE TYPE: peptide
347
--> 348 (ix) (iii) FEATURE INFORMATION:
349 (D) OTHER INFO: Xaa in position 2 is Met, Ile, Leu or Val;
350 Xaa in position 4 is Met, Ile, Leu or Val;
351 Xaa in position 6 is Asn, Asp, Gln or Glu.
352
353 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:
354
355 Thr Xaa Lys Xaa Arg Xaa
356 1 5
357
358

359 (2) INFORMATION FOR SEQ ID NO:20:
360
361 (i) SEQUENCE CHARACTERISTICS:
362 (A) LENGTH: 7 amino acids
363 (B) TYPE: amino acid
364 (C) STRANDEDNESS: single
365 (D) TOPOLOGY: linear
366
367 (ii) MOLECULE TYPE: peptide
368
--> 369 (ix) (iii) FEATURE INFORMATION:
370 (D) OTHER INFO: Xaa in position 1 is Met, Ile, Leu or Val;
371 Xaa in position 3 is Met, Ile, Leu or Val;
372 Xaa in position 5 is Met, Ile, Leu or Val;
373 Xaa in position 7 is Asn, Asp, Gln or Glu.
374
375 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:
376
377 Xaa Thr Xaa Lys Xaa Arg Xaa
378 1 5
379
380

381 (2) INFORMATION FOR SEQ ID NO:21:

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382
383 (i) SEQUENCE CHARACTERISTICS:
384 (A) LENGTH: 8 amino acids
385 (B) TYPE: amino acid
386 (C) STRANDEDNESS: single
387 (D) TOPOLOGY: linear
388
389 (ii) MOLECULE TYPE: peptide
390
--> 391 (ix) ~~(iii) FEATURE INFORMATION:~~
392 (D) OTHER ~~Info~~ Xaa in position 1 is Tyr or Phe;
393 Xaa in position 2 is Met, Ile, Leu or Val;
394 Xaa in position 4 is Met, Ile, Leu or Val;
395 Xaa in position 6 is Met, Ile, Leu or Val;
396 Xaa in position 8 is Asn, Asp, Gln or Glu.
397
398 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:
399
400 Xaa Xaa Thr Xaa Lys Xaa Arg Xaa
401 1 5
402
403
404
405

406 (2) INFORMATION FOR SEQ ID NO:22:
407
408 (i) SEQUENCE CHARACTERISTICS:
409 (A) LENGTH: 9 amino acids
410 (B) TYPE: amino acid
411 (C) STRANDEDNESS: single
412 (D) TOPOLOGY: linear
413
414 (ii) MOLECULE TYPE: peptide
415
--> 416 (iii) FEATURE INFORMATION:
417 Xaa in position 1 is Ala or Gly;
418 Xaa in position 2 is Tyr or Phe;
419 *same* Xaa in position 3 is Met, Ile, Leu or Val;
420 *even* Xaa in position 5 is Met, Ile, Leu or Val;
421 Xaa in position 7 is Met, Ile, Leu or Val;
422 Xaa in position 9 is Asn, Asp, Gln or Glu.
423
424 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:
425
426 Xaa Xaa Xaa Thr Xaa Lys Xaa Arg Xaa
427 1 5
428
429

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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/09/101,825

DATE: 01/14/2000
TIME: 00:09:21

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Line	Error	Original Text
348	Unknown or Misplaced Identifier	(iii) FEATURE INFORMATION:
369	Unknown or Misplaced Identifier	(iii) FEATURE INFORMATION:
391	Unknown or Misplaced Identifier	(iii) FEATURE INFORMATION:
416	Unknown or Misplaced Identifier	(iii) FEATURE INFORMATION: